



NOV 21 2002

PATENT &amp; TRADEMARK OFFICE

ATTORNEY DOCKET NO. 14114.0325U2

RECEIVED

NOV 29 2002

TECH CENTER 1600/2900

## SEQUENCE LISTING

*Sub C*

<110> Rosely M. Zancope-Oliveira  
Timothy J. Lott  
Leonard W. Mayer  
Errol Reiss  
George S. Deepe

<120> NUCLEIC ACIDS OF THE M ANTIGEN GENE OF  
HISTOPLASMA CAPSULATUM, ANTIGENS, VACCINES, AND ANTIBODIES,  
METHODS AND KITS FOR DETECTING HISTOPLASMOSES

<130> 14114.0325U2

<140> 09/674,195  
<141> 2000-10-26

<150> PCT/US99/09151  
<151> 1999-04-27

<150> 60/083,676  
<151> 1998-04-30

<160> 13

<170> FastSEQ for Windows Version 4.0

<210> 1  
<211> 3862  
<212> DNA  
<213> Histoplasma capsulatum

<220>  
<221> misc\_feature  
<222> 3258  
<223> n = g, a, c or t(u)

<400> 1

ggatcctgct	ggctccgata	actttgcttt	atccaagggt	ctcggcgaat	gccaggtgcc	60
atcgatctat	attttgaagt	ttatcaccc	aatggcttc	ccccatgacg	caccttttat	120
ttttatttc	attcatcttc	tctgtggcaa	acatgcaggt	atgcgagctc	tggaccctgg	180
ggtgtggccc	ttgatgcata	tggtttattt	atagccccc	ggaagccctg	gcctgttaaa	240
ttttggacct	cctcccgcca	ttctttccaa	acttcgtgcg	tccgtttccc	attttcccccc	300
tccccatttg	ggttccctat	aggccactgc	gtgctccact	caagaagggt	cccaagtcaat	360
ttggtcccta	ccctctccaa	cactatctgc	atatgtaata	tatatcgata	tctaaactgcc	420
attgattatt	tgtcttcttc	agcatctttt	tgtctcgagc	aagcttactc	cacgttcaat	480
tcagggggta	aaaatgcggt	cgctcaagct	tatactcgcc	tccgcgggtg	ttgtttctgc	540
agcctgtccc	tacatgtcag	gggagatgcc	tagcggtcag	aaaggcccc	tcgatcgccg	600
ccatgacact	ctctccgacc	ctacggacca	gtttcttagc	aagttttaca	ttgacgatga	660
acagtcggtg	ctaacaacgg	acgtgggtgg	tcccatcgag	gaccaacaca	gcctgaaggc	720
tggaaataga	ggcccaactc	tacttgagga	ttttatcttc	cgccagaaga	ttcaacactt	780
tgatcatgag	agggtatgta	gataaaaaat	atgtgaccgt	gttgcaaaatc	cgctaattca	840
attttacgca	ggttcctgag	cgcgcccgtcc	atgctcgagg	agctggtgcc	catggcgtat	900

tcacatccta taataactgg tcgaatatca cagccgcata cttcttgaac gcggcaggaa  
agcagacacc agtattcgta cggtttcta cagtcgctgg tagcagaggc agtggact  
ctgctcgca tatccacgg a ttgcgaccc g tctgtatac cgatgaaggc aattttgta  
agcattatat cgtggtagtc atactataa cagcacaaca aatataaataa caaaccagg  
acctaggctg actactcgcc aatgttagata tcgctggaaa caacgttcca gtcttctca  
ttcaggacgc tattcaattt cctgatttga ttcacgctgt caagcccaa ccagacagt  
aaattcccc ggtgcaact gcacatgata cggcatggg tttctcagc cagcagccca  
gctcatgca tccccctttt tggcaatgt caggacatgg aatccctcg tcaatgcgtc  
atgtttaggg gttttttttt catacccttc gacttgcac cgacgaggc aactcgac  
tggtaagtt tcgctggaa accctccaag gaagagcggg cctggatgg gaagaggcac  
aggctttgg cggaaagaat cccgacttcc atcgacaaga cctctggat gccattgaat  
ctggaaaggta cccctgaggta gaggttaagat atgattcccc caaatcatta gttctgac  
tggttctctg ctctgtcggt tgcttttttgc ttttttttgc atatcttcaa ctaagactga  
ctttatatac gtttactca tatagtctggg ctttcaatttgc tgaatgaag cagatcaatc  
caagttgtat ttcatgtat tagatccccac caaatcatttgc ccagaagaac ttgttctt  
cccccaatc gggaaaatgg ttttgcaccg aaacccaaaa agttattttgc cggaaactga  
gcagatcatg gttggccac cccctatata ttttgcatttgc gaatacatgt atagctagat  
gaagcgata tctaaatata tttccacagt tccaccagg tcatgttagt cgccgaaatcg  
atttcacggta tgacccttttgc cttcaggggcc gtttgcactc ctaccttgc actcaatttga  
atcgccatgg aggtcccaac ttgcagcaac tgccgatcaaa cagaccccg atcccttcc  
ataacaacaa tcgcaacggt gctggtaagc tacttctcac ctaccatgtc aacttccatc  
ttgacccaaat cgtttgtat agagtattaa catccccgtc tgcacaggac aaatgttcat  
ccctctaaac acggccgcat atacacccaa ctcaatgac aacggattcc cacaacaagg  
caacccggacc cataacacagg gattttcac cgcacccctggg cgtatggtaa atggaccact  
agtgcgcgag ctccggccca gcttcaacga cgtctggcc caaccgcgtc tcttctacaa  
ctcaactacg gtcttcgaga agcaatttgc cgtcaacgc atgcgttgc aaaactcccc  
cgtgcggagt gaaaccgtgc gtaagaacgt catcatccag ctgaaccgcg tcgacaacga  
ctctggccgc cgcgtcgcc tagctatcg gtcgaaaccc ccattcccg acccaaccc  
ctaccacaac aaggcaaccc tccccatcgg cacccctggc acgaatctcc tgcggctcga  
cgggctgaaa atcgccctcc tgacaagaga cgacggtagc ttacacgtcg cggagcagct  
ccggcccgcc ttaacacggc ccaacacaa agtagatatac gtcctagtgg gctcatcgct  
tgatccccaa cgcggcgtga acatgaccta ttccggcgcc gacggctga tcttcgatgc  
cgtgatcgta gtcggccggcc tgctcacgg cgcctcaacg caatacccaa gaggtcgccc  
gctcaggatt attacggatg catacgcgtt gggaaagccc gttggcccg tcggtgacgg  
tagcaatgaa gccttcgttgc acgtccttgc ggcgcgttgc gggatgcgt cgaatggct  
ggaccagccc ggtgtgtata ttccaaacga tggatgttag gcttacgttga gaaatgtctt  
ggacggattt acggcatatc gtttttttgc gtcgttcccg ttggatagaa gcttggatgg  
aggttttggg cggaaatatg gtttactac cccccccccc ccctttttt ttttccctt  
ctgtttttcc atctttgggtt gaggttaat tgcagatatac agttaatttgc gtttacaaa  
gccggtgtca agtttcanga ggcctaatta atttgaagag gaggttgaag tgaaatcttgc  
gtgttaactat aataattttt aataactat aacttataat taatgtctat tgtaatttcc  
tctcacattt aatctatatt tgatccttgc ctttgcgttgc tgtttaataa taagccaaaga  
gagacaaata atgatagatt aacaaataat tgcacacccca ataggccctc cctcagata  
tcagatatta tctatcatgt tgtaatgtataa cctcaaaaat gccacaagct tgcctgatata  
tgaatatttta tatgtgttaa atgttagggaa gagcgtacca tccaaataac cagaaaaac  
tgtttttagct taaaatctca ctaaggtcggt tcgtgtctat ttggaaatggc tgccggcaagc  
tgactatctg ataaaaatgt ctgtatttcc gtttgcacggc gcatgttgcgactttca  
atagataaaa cctgaacacgt ttagcccttgc ttggggggaa taggggttag gggggccgg  
tacatatcat tccatataat cccaaacta aaatagatatac atatataat atatataat  
acaacacccctt caaaaaaggat cc

<210> 2  
<211> 707  
<212> PRT  
<213> *Histoplasma capsulatum*

<400> 2  
Met Pro Ser Gly Gln Lys Gly Pro Leu Asp Arg Arg His Asp Thr Leu  
1 5 10 15  
Ser Asp Pro Thr Asp Gln Phe Leu Ser Lys Phe Tyr Ile Asp Asp Glu  
20 25 30  
Gln Ser Val Leu Thr Thr Asp Val Gly Gly Pro Ile Glu Asp Gln His  
35 40 45  
Ser Leu Lys Ala Gly Asn Arg Gly Pro Thr Leu Leu Glu Asp Phe Ile  
50 55 60  
Phe Arg Gln Lys Ile Gln His Phe Asp His Glu Arg Val Pro Glu Arg  
65 70 75 80  
Ala Val His Ala Arg Gly Ala Gly His Gly Val Phe Thr Ser Tyr  
85 90 95  
Asn Asn Trp Ser Asn Ile Thr Ala Ala Ser Phe Leu Asn Ala Ala Gly  
100 105 110  
Lys Gln Thr Pro Val Phe Val Arg Phe Ser Thr Val Ala Gly Ser Arg  
115 120 125  
Gly Ser Val Asp Ser Ala Arg Asp Ile His Gly Phe Ala Thr Arg Leu  
130 135 140  
Tyr Thr Asp Glu Gly Asn Phe Asp Ile Val Gly Asn Asn Val Pro Val  
145 150 155 160  
Phe Phe Ile Gln Asp Ala Ile Gln Phe Pro Asp Leu Ile His Ala Val  
165 170 175  
Lys Pro Gln Pro Asp Ser Glu Ile Pro Gln Ala Ala Thr Ala His Asp  
180 185 190  
Thr Ala Trp Asp Phe Leu Ser Gln Gln Pro Ser Ser Leu His Ala Leu  
195 200 205  
Phe Trp Ala Met Ser Gly His Gly Ile Pro Arg Ser Met Arg His Val  
210 215 220  
Asp Gly Trp Gly Val His Thr Phe Arg Leu Val Thr Asp Glu Gly Asn  
225 230 235 240  
Ser Thr Leu Val Lys Phe Arg Trp Lys Thr Leu Gln Gly Arg Ala Gly  
245 250 255  
Leu Val Trp Glu Glu Ala Gln Ala Leu Gly Gly Lys Asn Pro Asp Phe  
260 265 270  
His Arg Gln Asp Leu Trp Asp Ala Ile Glu Ser Gly Arg Tyr Pro Glu  
275 280 285  
Trp Glu Leu Gly Phe Gln Leu Val Asn Glu Ala Asp Gln Ser Lys Phe  
290 295 300  
Asp Phe Asp Leu Leu Asp Pro Thr Lys Ile Ile Pro Glu Glu Leu Val  
305 310 315 320  
Pro Phe Thr Pro Ile Gly Lys Met Val Leu Asn Arg Asn Pro Lys Ser  
325 330 335  
Tyr Phe Ala Glu Thr Glu Gln Ile Met Phe Gln Pro Gly His Val Val  
340 345 350  
Arg Gly Ile Asp Phe Thr Asp Asp Pro Leu Leu Gln Gly Arg Leu Tyr  
355 360 365

Ser Tyr Leu Asp Thr Gln Leu Asn Arg His Gly Gly Pro Asn Phe Glu  
370 375 380  
Gln Leu Pro Ile Asn Arg Pro Arg Ile Pro Phe His Asn Asn Asn Arg  
385 390 395 400  
Asp Gly Ala Gly Gln Met Phe Ile Pro Leu Asn Thr Ala Ala Tyr Thr  
405 410 415  
Pro Asn Ser Met Ser Asn Gly Phe Pro Gln Gln Ala Asn Arg Thr His  
420 425 430  
Asn Arg Gly Phe Phe Thr Ala Pro Gly Arg Met Val Asn Gly Pro Leu  
435 440 445  
Val Arg Glu Leu Ser Pro Ser Phe Asn Asp Val Trp Ser Gln Pro Arg  
450 455 460  
Leu Phe Tyr Asn Ser Leu Thr Val Phe Glu Lys Gln Phe Leu Val Asn  
465 470 475 480  
Ala Met Arg Phe Glu Asn Ser His Val Arg Ser Glu Thr Val Arg Lys  
485 490 495  
Asn Val Ile Ile Gln Leu Asn Arg Val Asp Asn Asp Leu Ala Arg Arg  
500 505 510  
Val Ala Leu Ala Ile Gly Val Glu Pro Pro Ser Pro Asp Pro Thr Phe  
515 520 525  
Tyr His Asn Lys Ala Thr Val Pro Ile Gly Thr Phe Gly Thr Asn Leu  
530 535 540  
Leu Arg Leu Asp Gly Leu Lys Ile Ala Leu Leu Thr Arg Asp Asp Gly  
545 550 555 560  
Ser Phe Thr Ile Ala Glu Gln Leu Arg Ala Ala Phe Asn Ser Ala Asn  
565 570 575  
Asn Lys Val Asp Ile Val Leu Val Gly Ser Ser Leu Asp Pro Gln Arg  
580 585 590  
Gly Val Asn Met Thr Tyr Ser Gly Ala Asp Gly Ser Ile Phe Asp Ala  
595 600 605  
Val Ile Val Val Gly Gly Leu Leu Thr Ser Ala Ser Thr Gln Tyr Pro  
610 615 620  
Arg Gly Arg Pro Leu Arg Ile Ile Thr Asp Ala Tyr Ala Tyr Gly Lys  
625 630 635 640  
Pro Val Gly Ala Val Gly Asp Gly Ser Asn Glu Ala Leu Arg Asp Val  
645 650 655  
Leu Met Ala Ala Gly Gly Asp Ala Ser Asn Gly Leu Asp Gln Pro Gly  
660 665 670  
Val Tyr Ile Ser Asn Asp Val Ser Glu Ala Tyr Val Arg Ser Val Leu  
675 680 685  
Asp Gly Leu Thr Ala Tyr Arg Phe Leu Asn Arg Phe Pro Leu Asp Arg  
690 695 700  
Ser Leu Val  
705

<210> 3  
<211> 8  
<212> PRT  
<213> *Histoplasma capsulatum*

<400> 3  
Ser Asp Pro Thr Asp Gln Phe Leu  
1 5

<210> 4  
<211> 15  
<212> PRT  
<213> *Histoplasma capsulatum*

<400> 4  
Asp Phe Ile Phe Arg Gln Lys Ile Gln His Phe Asp His Glu Arg  
1 5 10 15

*B1*  
<210> 5  
<211> 9  
<212> PRT  
<213> *Histoplasma capsulatum*

<400> 5  
Thr Leu Gln Gly Arg Ala Gly Leu Val  
1 5

<210> 6  
<211> 16  
<212> PRT  
<213> *Histoplasma capsulatum*

<400> 6  
Ala Gln Ala Leu Gly Gly Lys Asn Pro Asp Phe His Arg Gln Asp Leu  
1 5 10 15

<210> 7  
<211> 6  
<212> PRT  
<213> *Histoplasma capsulatum*

<400> 7  
Ser Gly Arg Tyr Pro Glu  
1 5

<210> 8  
<211> 10  
<212> PRT  
<213> Histoplasma capsulatum

<400> 8  
Phe Asp Phe Asp Leu Leu Asp Pro Thr Lys  
1 5 10

<210> 9  
<211> 14  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence; M antigen-specific oligonucleotide

<400> 9  
Ile Ile Pro Glu Glu Leu Val Pro Phe Thr Pro Ile Gly Lys  
1 5 10

<210> 10  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence; sense amplification primer

<220>  
<221> misc\_feature  
<222> 3  
<223> r = a or g

<220>  
<221> misc\_feature  
<222> 6, 12, 15  
<223> y = c or t

<220>  
<221> misc\_feature  
<222> 9  
<223> v = g, c or a

<400> 10  
aaraayccvg aytty

<210> 11  
<211> 14  
<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence; anti-sense amplification primer

<220>

<221> misc\_feature

<222> 3, 9

<223> n = g, a, c or t(u)

<220>

<221> misc\_feature

<222> 6

<223> d = g or a

<220>

<221> misc\_feature

<222> 12

<223> r = a or g

<400> 11

tttnccdatng traa

14

<210> 12

<211> 22

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence; sense cDNA amplification primer

<400> 12

cggaatcctc cgaccctacg ga

22

<210> 13

<211> 27

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence; anti-sense cDNA amplification primer

<400> 13

accaagcttc tatccaacgg gaaccgaa

27